

Figure 1

Sequence alignment of mouse Serca 1, 2 and 3 protein.

5	Serca1a	1	MEAAHSKSTEECLSYFGVSETTGLTPDQVKRHLEKYGNELPAEEGKSLWELVVEQFEDL
	Serca2a	1	..N..T.TV..V.GH...N.S...SLE...KLK.RW.S.....T.L...I.....
	Serca2b	1	..N..T.TV..V.GH...N.S...SLE...KLK.RW.S.....T.L...I.....
	Serca3a	1	..E..LL.AADV.RR.S.TAEG..SLE..TDAR.R.....T.....
	Serca3b	1	..E..LL.AADV.RR.S.TAEG..SLE..TDAR.R.....T.....
10	Serca3c	1	..E..LL.AADV.RR.S.TAEG..SLE..TDAR.R.....T.....
	Serca1a	61	LVRILLLAACISFVLAWFEEGEETVTAFFVEPFVILLILIANAIVGVWQERNAENAIEALK
	Serca2a	61	.....I.....V.....
	Serca2b	61	.....I.....V.....
15	Serca3a	61	.....LV.....T.....L..M..V.....S.....
	Serca3b	61	.....LV.....T.....L..M..V.....S.....
	Serca3c	61	.....LV.....T.....L..M..V.....S.....
	Serca1a	121	EYEPENMGKVYRADRKSVQRIKARDIVPGDIVEVAVGDKVPADIRILSIKSTTLRVDQSIL
20	Serca2a	121	.....Q.....K.....I.....LT.....
	Serca2b	121	.....Q.....K.....I.....LT.....
	Serca3a	121	.....I.S...G...R.....L.LIE.....
	Serca3b	121	.....I.S...G...R.....L.LIE.....
	Serca3c	121	.....I.S...G...R.....L.LIE.....
25	Serca1a	181	TGESVSVIKHTDPVPDPRAVNQDKKNMLFSGTNIAAGKAVGIVATTGVSTEIGKIRDQMA
	Serca2a	181	.....M.V.VA...N.....E.V
	Serca2b	181	.....M.V.VA...N.....E.V
	Serca3a	181	.....T...AI.....S...L.VAVA..LQ..L...S...
30	Serca3b	181	.....T...AI.....S...L.VAVA..LQ..L...S...
	Serca3c	181	.....T...AI.....S...L.VAVA..LQ..L...S...
	Serca1a	241	ATEQDKTPLQOKLDEFGEQLSKVISLICVAVWLINIGHFNDPVHGGSWFRGAIYYFKIAV
35	Serca2a	241	....ER.....I...I.....I.....
	Serca2b	241	....ER.....I...I.....I.....
	Serca3a	241	.V.PER...R.....R..HA..V.....V.....A..A....L..V.....
	Serca3b	241	.V.PER...R.....R..HA..V.....V.....A..A....L..V.....
	Serca3c	241	.V.PER...R.....R..HA..V.....V.....A..A....L..V.....
40	Serca1a	301	ALAVAAIPEGLPAVITTCLALGTRMAKNNAIVRSLPSVETLGCTSVICSDKTGTLTTNQ
	Serca2a	301	.....
	Serca2b	301	.....
	Serca3a	301	.....R.....
	Serca3b	301	.....R.....
45	Serca3c	301	.....R.....
	Serca1a	361	MSVCKMFIIDKVDGDCSLNEFSITGSTYAPEGEVLKNDKPVVRAGQYDGLVELATICALC
	Serca2a	361	....R...L...E..T.....I...Q.D...KCH.....
	Serca2b	361	....R...L...E..T.....I...Q.D...KCH.....
50	Serca3a	361	....R...VVAEAEAGT.R.H..T.S.T..T....RQGEQ...C..F.....
	Serca3b	361	....R...VVAEAEAGT.R.H..T.S.T..T....RQGEQ...C..F.....
	Serca3c	361	....R...VVAEAEAGT.R.H..T.S.T..T....RQGEQ...C..F.....
55	Serca1a	421	NDSSSLDFNETKGVYEKVGATETALTTLVEKMNVFNTVRSLSKVERANACNSVIRQLMK
	Serca2a	421	...A..Y..A.....C.....D..LKG...I.....K....
	Serca2b	421	...A..Y..A.....C.....D..LKG...I.....K....
	Serca3a	421	...A..Y..A.....C.....D..DLKG...R....G.....K...R
	Serca3b	421	...A..Y..A.....C.....D..DLKG...R....G.....K...R
	Serca3c	421	...A..Y..A.....C.....D..DLKG...R....G.....K...R

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	Serca1a	481	KEFTLEFSRDRKSMSVYCSPAKSSRAAVGNKMFVKGAPEGVIDRCNYVRVGTTTRVPLTGP
	Serca2a	481	.....T.N.P..TSMS-.....THI...S.K..M.PG
	Serca2b	481	.....T.N.P..TSMS-.....THI...S.K..M.PG
	Serca3a	481	.....T.TRADPKVQ.S.....S..E..SS....SRTA..STT
5	Serca3b	481	.....T.TRADPKVQ.S.....S..E..SS....SRTA..STT
	Serca3c	481	.....T.TRADPKVQ.S.....S..E..SS....SRTA..STT
	Serca1a	541	VKEKIMSVIKEWGTGRDTRLCLALATRDTPPKREEMVLDDSAKFMEYEMDLTFVGVVGML
	Serca2a	540	..Q.....R...S.S.....H.N.L.....H.E...N.IK..TN....C....
10	Serca2b	540	..Q.....R...S.S.....H.N.L.....H.E...N.IK..TN....C....
	Serca3a	541	SR.H.LAK.RD..S.S.....RK.D.H...CSR.VQ..T.....C....
	Serca3b	541	SR.H.LAK.RD..S.S.....RK.D.H...CSR.VQ..T.....C....
	Serca3c	541	SR.H.LAK.RD..S.S.....RK.D.H...CSR.VQ..T.....C....
15	Serca1a	601	DPPRKEVTGSIQLCRDAGIRVIMITGDNKGTAIAICRRIGIFSENEEVTDRAYTGREFDD
	Serca2a	600	....I..AS.VK...Q.....V.....GQD.D..SK.F.....E
	Serca2b	600	....I..AS.VK...Q.....V.....GQD.D..SK.F.....E
	Serca3a	601	....P..AAC.TR.SR....V.....V.....L...GDT.D.LGK.....
	Serca3b	601	....P..AAC.TR.SR....V.....V.....L...GDT.D.LGK.....
20	Serca3c	601	....P..AAC.TR.SR....V.....V.....L...GDT.D.LGK.....
	Serca1a	661	LPLAEQREACRRACCFARVEPSHKSKIIVEYLQSYDEITAMTGDGVNDAPALKKAEIGIAM
	Serca2a	660	.SPSA..D..LN.R.....F..F.....S.....
	Serca2b	660	.SPSA..D..LN.R.....F..F.....S.....
25	Serca3a	661	.SPEQ..Q...T.R.....A...R...N...FN.....
	Serca3b	661	.SPEQ..Q...T.R.....A...R...N...FN.....
	Serca3c	661	.SPEQ..Q...T.R.....A...R...N...FN.....
	Serca1a	721	GSGTAVAKTASEMVLADDNFSTIVAAVEEGRAIYNNMKQFIRYLISSNVGEVVCIFLTAA
30	Serca2a	720	.....
	Serca2b	720	.....
	Serca3a	721	.....S.A...S...AS.....I
	Serca3b	721	.....S.A...S...AS.....I
	Serca3c	721	.....S.A...S...AS.....I
35	Serca1a	781	LGLPEALIPVQLLWVNLVTDGLPATALGFNPPDLDIMDRPPRSPEPLISGWLFFRYMAI
	Serca2a	780	..F.....NK...N.....L..
	Serca2b	780	..F.....NK...N.....L..
	Serca3a	781	.....EK...N.R.A.....L..
40	Serca3b	781	.....EK...N.R.A.....L..
	Serca3c	781	.....EK...N.R.A.....L..
	Serca1a	841	GGYVGAATVGAAAWFLYAEDGPHVSYHQLTHFMQCTEHNPEFDGLDCEVFPEPEPMTMA
	Serca2a	840	.C.....IA.DG..R..FY..S..L..K.D..D...V..AI..S.Y.....
45	Serca2b	840	.C.....IA.DG..R..FY..S..L..K.D..D...V..AI..S.Y.....
	Serca3a	841	.V...L...A..T....DAE..Q.TFY..RN.LK.S.D..L.A.I..K...SRF.T...
	Serca3b	841	.V...L...A..T....DAE..Q.TFY..RN.LK.S.D..L.A.I..K...SRF.T...
	Serca3c	841	.V...L...A..T....DAE..Q.TFY..RN.LK.S.D..L.A.I..K...SRF.T...
50	Serca1a	901	LSVLVTIEMCNALNSLSENQSLLRMPWVNIWLLGSICLSMSLHFLILYVDPLPMIFKLR
	Serca2a	900	.....E.....V.....E...L..QIT
	Serca2b	900	.....E.....V.....E...L..QIT
	Serca3a	901	.....V.....L.P....AVVM..A.....L.P...L..QVT
	Serca3b	901	.....V.....L.P....AVVM..A.....L.P...L..QVT
55	Serca3c	901	.....V.....L.P....AVVM..A.....L.P...L..QVT

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Serca1a 961 ALDFTQWLMVLKISLPVIGLDELLKFIARNYLEG  
Serca2a 960 P.NL.....LM..T...V.....QPATLE  
Serca2b 960 P.NL.....LM..T...V.....QPGKECVQPATKSSCSLSACTDGISWP  
Serca3a 961 P.SGR..GV..QM.....L...A..YLS..HMDEKKDLK  
5 Serca3b 961 P.SGR..GV..QM.....L...A..YLS..HMD.VLGTFMQARSROLPTTSRTPYHTGKK  
Serca3c 961 P.SGR..GV..QM.....L...A..YLS..HMD.VLGTFMQARSROLPTTSRTPYHTGLA  
  
Serca2b 1020 FVLLIMPLVVVVYSTDTNFSDFMWS  
10 Serca3b 1021 GPEVNPGSRGESPVWPSD  
Serca3c 1021 SWKKRT

Figure 2

Sequence similarity of Serca2 proteins in mammalian species

	Mouse_2a	1	MENAHTKTVEEVLGHFGVNESTGLSLEQVKLKERWGSNELPAEEGKTLLLELVIEQFEDL
5	Mouse_2b	1	.....
	Rat_2b	1	.....
	Rat_2a	1	.....
	Dog_2a	1	.....
	Cat_2a	1	.....Y.....
10	Pig_2a	1	.....
	Pig_2b	1	.....
	Human_2b	1	.....
	Human_2c	1	.....
	Human_2a	1	.....
15	Rabbit_2a	1	.....
	Rabbit_2b	1	.....
	Mouse_2a	61	LVRILLLAACISFVLAWFEEGEETITAFVEPFVILLILVANAIVGVWQERNAENAIEALK
	Mouse_2b	61	.....
20	Rat_2b	61	.....
	Rat_2a	61	.....
	Dog_2a	61	.....
	Cat_2a	61	.....
	Pig_2a	61	.....
25	Pig_2b	61	.....
	Human_2b	61	.....
	Human_2c	61	.....
	Human_2a	61	.....
	Rabbit_2a	61	.....
30	Rabbit_2b	61	.....
	Mouse_2a	121	EYEPENMGKVYRQDRKSVQRIKAKDIVPGDIVEIAVGDKVPADIRLTSIKSTTLRVDQSIL
	Mouse_2b	121	.....
	Rat_2b	121	.....
35	Rat_2a	121	.....
	Dog_2a	121	.....
	Cat_2a	121	.....
	Pig_2a	121	.....
	Pig_2b	121	.....
40	Human_2b	121	.....
	Human_2c	121	.....
	Human_2a	121	.....
	Rabbit_2a	121	.....
	Rabbit_2b	121	.....
45	Mouse_2a	181	TGESVSVIKHTDPVPDPRAVNQDKKNMLFSGTNIAAGKAMGVVVATGVNTEIGKIRDEM
	Mouse_2b	181	.....
	Rat_2b	181	.....
	Rat_2a	181	.....
50	Dog_2a	181	.....
	Cat_2a	181	.....
	Pig_2a	181	.....
	Pig_2b	181	.....
	Human_2b	181	.....
55	Human_2c	181	.....
	Human_2a	181	.....
	Rabbit_2a	181	.....
	Rabbit_2b	181	.....

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	Mouse_2a	241	ATEQERTPLQQKLDEFGEQLSKVISLICIAVWIINIGHFNDFVHGGSWIRGAIYYFKIAV
	Mouse_2b	241	.....
	Rat_2b	241	.....
5	Rat_2a	241	.....
	Dog_2a	241	.....
	Cat_2a	241	.....
	Pig_2a	241	.....
	Pig_2b	241	.....
10	Human_2b	241	.....
	Human_2c	241	.....
	Human_2a	241	.....
	Rabbit_2a	241	.....
	Rabbit_2b	241	.....
15	Mouse_2a	301	ALAVAAIPEGLPAVITTCLALGTRRMAKKNAIVRSLPSVETLGCTSVICSDKTGTLTTNQ
	Mouse_2b	301	.....
	Rat_2b	301	.....
	Rat_2a	301	.....
20	Dog_2a	301	.....
	Cat_2a	301	.....
	Pig_2a	301	.....
	Pig_2b	301	.....
	Human_2b	301	.....
25	Human_2c	301	.....
	Human_2a	301	.....
	Rabbit_2a	301	.....
	Rabbit_2b	301	.....
30	Mouse_2a	361	MSVCRMFIELDKVEGDTCSLNEFSITGSTYAPIGEVQKDDKPKVCHQYDGLVELATICALC
	Mouse_2b	361	.....
	Rat_2b	361	.....T.....
	Rat_2a	361	.....T.....
	Dog_2a	361	.....R.....S.....T.....H.....
35	Cat_2a	361	.....T.....H.....
	Pig_2a	361	.....T.....H.....
	Pig_2b	361	.....T.....H.....
	Human_2b	361	.....R.....T.....H.....N.....
	Human_2c	361	.....R.....T.....H.....N.....
40	Human_2a	361	.....R.....T.....H.....N.....
	Rabbit_2a	361	.....D.....T.....H.....
	Rabbit_2b	361	.....D.....T.....H.....
45	Mouse_2a	421	NDSALDYNEAKGVYEKVGEATETALTCLVEKMNVFDTLKGLSKIERANACNSVIKQLMK
	Mouse_2b	421	.....
	Rat_2b	421	.....
	Rat_2a	421	.....
	Dog_2a	421	.....
	Cat_2a	421	.....K.F.....
50	Pig_2a	421	.....
	Pig_2b	421	.....
	Human_2b	421	.....
	Human_2c	421	.....
	Human_2a	421	.....
55	Rabbit_2a	421	.....
	Rabbit_2b	421	.....

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	Mouse_2a	481	KEFTLEFSRDRKSMSVYCTPNKPSRTSMSKMFVKGAPEGVIDRCTHIRVGSTKVPMTPGV
	Mouse_2b	481	.....
	Rat_2b	481	.....
	Rat_2a	481	.....
5	Dog_2a	481	.....
	Cat_2a	481	.....
	Pig_2a	481	.....
	Pig_2b	481	.....
	Human_2b	481	.....S..
10	Human_2c	481	.....S..
	Human_2a	481	.....S..
	Rabbit_2a	481	.....A..
	Rabbit_2b	481	.....A..
15	Mouse_2a	541	KQKIMSVIREWGSGSDTLRCLALATHDNPLKREEMHLEDSANFIKYETNLITFVGCVGMLD
	Mouse_2b	541	.....
	Rat_2b	541	.....R..
	Rat_2a	541	.....R..
	Dog_2a	541	...V.....R...N.....
20	Cat_2a	541	...V.....R...N.....
	Pig_2a	541	.....MR...N.....
	Pig_2b	541	.....MR...N.....
	Human_2b	541	.....R.....
	Human_2c	541	.....R.....
25	Human_2a	541	.....R.....
	Rabbit_2a	541	.....R.....K.....
	Rabbit_2b	541	.....R.....K.....
30	Mouse_2a	601	PPRIEVASSVKLCRQAGIRVIMITGDNKGTAVAICRRIGIFGQDEDVTSKAFTGREFDEL
	Mouse_2b	601	.....
	Rat_2b	601	.....
	Rat_2a	601	.....
	Dog_2a	601	.....
	Cat_2a	601	.....
35	Pig_2a	601	.....
	Pig_2b	601	.....
	Human_2b	601	.....
	Human_2c	601	.....
	Human_2a	601	.....
40	Rabbit_2a	601	.....E...A.....
	Rabbit_2b	601	.....E...A.....
45	Mouse_2a	661	SPSAQRDACLNARCFARVEPSHKSKIVEFLQSFDEITAMTGDGVNDAPALKKSEIGIAMG
	Mouse_2b	661	.....
	Rat_2b	661	.....
	Rat_2a	661	.....
	Dog_2a	661	.....
	Cat_2a	661	.....
	Pig_2a	661	N.....E.....
50	Pig_2b	661	N.....E.....
	Human_2b	661	N.....A.....
	Human_2c	661	N.....A.....
	Human_2a	661	N.....A.....
	Rabbit_2a	661	N.....A.....
55	Rabbit_2b	661	N.....A.....

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	Mouse_2a	721	SGTAVAKTASEMVLADDNFSTIVAAVEEGRAIYNNMKQFIRYLISSNVGEVVCIFLTAAL
	Mouse_2b	721	.....
	Rat_2b	721	.....
	Rat_2a	721	.....
5	Dog_2a	721	.....
	Cat_2a	721	.....
	Pig_2a	721	.....
	Pig_2b	721	.....
	Human_2b	721	.....
10	Human_2c	721	.....
	Human_2a	721	.....
	Rabbit_2a	721	.....
	Rabbit_2b	721	.....
15	Mouse_2a	781	GFPEALIPVQLLWVNLVTDGLPATALGFNPPDLDIMNKPFRNPKEPLISGWLFFRYLAIG
	Mouse_2b	781	.....
	Rat_2b	781	.....
	Rat_2a	781	.....
	Dog_2a	781	.....
20	Cat_2a	781	.....
	Pig_2a	781	.....
	Pig_2b	781	.....
	Human_2b	781	.....
	Human_2c	781	.....
25	Human_2a	781	.....
	Rabbit_2a	781	.....
	Rabbit_2b	781	.....
30	Mouse_2a	841	CYVGAATVGAAAWWFIAADGGPRVSFYQLSHFLQCKEDNPDFDGVDCAI FESPYPMTMAL
	Mouse_2b	841	.....
	Rat_2b	841	.....E.....
	Rat_2a	841	.....E.....
	Dog_2a	841	.....D.....E.....
	Cat_2a	841	.....D.....E.....
35	Pig_2a	841	.....T.....E.....V.....
	Pig_2b	841	.....T.....E.....V.....
	Human_2b	841	.....E.....
	Human_2c	841	.....E.....
	Human_2a	841	.....E.....
40	Rabbit_2a	841	.....E.....
	Rabbit_2b	841	.....E.....
45	Mouse_2a	901	SVLVTIEMCNALNSLSENQSLLRMPWENIWLVGSI CLSMLHFLILYVEPLPLIFQITP
	Mouse_2b	901	.....
	Rat_2b	901	.....
	Rat_2a	901	.....
	Dog_2a	901	.....
	Cat_2a	901	.....
	Pig_2a	901	.....
50	Pig_2b	901	.....
	Human_2b	901	.....
	Human_2c	901	.....
	Human_2a	901	.....
	Rabbit_2a	901	.....
55	Rabbit_2b	901	.....

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Mouse\_2a 961 LNL TQWLMVLKISLPVILMDET LKFVARNYLEQPAILE-----  
Mouse\_2b 961 .....GKECVQPATKSSCSLSACTDGISWPF  
Rat\_2b 961 .....GKECA.....P.....  
Rat\_2a 961 .....AILE  
5 Dog\_2a 961 .....AILE  
Cat\_2a 961 .....AILE  
Pig\_2a 961 .....AILE  
Pig\_2b 961 .....GKEC.....F.....  
Human\_2b 961 ..V.....GKEC.....F.....  
10 Human\_2c 961 ..V.....VLSSL  
Human\_2a 961 ..V.....AILE  
Rabbit\_2a 961 ..V.....AILE  
Rabbit\_2b 961 ..V.....GKEC....PQ....W....E.V....  
  
15 Mouse\_2b 1021 VLLIMPLVVVVYSTDTNFSDMEFS  
Rat\_2b 1020 .....  
Pig\_2b 1019 .....  
Human\_2b 1019 .....  
Rabbit\_2b 1019 ....V...M.....LL..  
20



**Fig. 3 Targeting construct for Serca2 flox gene modification. Sequence information.**

----- Serca2 gene -----

**LoxP site 1: Intron 1**  
underlined sequence = loxP site and cloning sequence

**Exon I (partial)**

5 CACGGGGCTGAGCTTGGAGCAGGTCAAGAAGCTCAAGGAGAGATGGGGCTCCAACGgtaggtgcggggcg  
ccggggcgtgcagcggcgcgggcgggcccgagcgccaaggaagatggctgacccggctccacccctcgtggg  
gcttggctcggcgcgggcgcccgacggtcgcgagagggccggcggtccacgcgcggtctgggccaatcgccg  
10 accttaggggtctcgaatcaagcttatcgataccgtcgatcggaacctcgagggggggccgggtacccggg  
gatcaattcgagctcgcccggggatcgatccggaaccttaataATAACTTCGTATAATGTATGCTATACG  
AAGTTATtaggtccctcgacctgcagcccaagctccGGGGAatctcgagccgggtgaccttcccgccggcg  
ctcagcgagtcggattgggggggggggagagggagtggaaggagggaggaggttcttcgcggtgggctg  
agtcccccgcgatttatgagggcgatggttggtagaaaccttcggacctttcttctgctccccaaa  
15 gttgcacatctggcagaagtgatgaccagctgaaatgactgcatggctcctggaggccggagagggctta  
cgggcagttccgagggccactgattaccagggctgaataattttctcggggtatcaaagtggagacagatt  
gttgtagcttcatacacctatatccgccattcagacaacgatgggtggtagaatttagcagtttttaataaa  
agcgctaataacaatatcttcattttttctttc

----- Serca2 gene -----

**LoxP site 2: Intron 3 5' of genomic XcmI site**  
underlined sequence = loxP site, cloning sites and partial HSV-TK

20 ccaatttttattcttagaacaattgtattcttatactgtgataggaagtgaataatcatacagtaacttgtc  
ttagggtttcacaaaactgataaactgtatgggttcaattatgtattcacacgtttaagtctgacccagggG  
GATCCggaaccttaataATAACTTCGTATAATGTATGCTATACGAAGTTATtaggtccctcgacctgcag  
25 cccaagctgatcccttagtcgagcccccagctgggttctttccgctcagaagccatagagcccccagaata  
cccagcatgcctgctattgtcttcccaatcctcccccttgctgtcctgccccacccccacccccagaata  
gaatgacacctactcagacaatgcgatgcaatttctctcattttatttaggaaaggacagtgaggagtggcac  
cttcacgggtcaaggaaggcacgggggaggggcaaacacagatggctggcaactagaaggcacagtcga  
ggctgatcagcgagctctagctagagaattgatccctcagaagaactcgtcaagaaggcgatagaaggc  
30 gatgcgctgcgaatcgggagcggcgat\*ccgtaaaagcacgaggaagcgg\*cgcccatctcgccgccaagc  
tctttcagcaatatcacgggtagccaacgctatgt\*ctgataagcgggtccgcccacacccaa\*ccggcca  
caagtc\*atgaaatcca\*aaaaagcggggccatttttccacc\*atgatttt\*cgqcaagcaaggccctt\*c  
cattgggtcaccgac\*aga\*catt\*tcggt\*c\*ggcattgcgc\*ccct

----- HSV-TK Neo antibiotics cassette -----

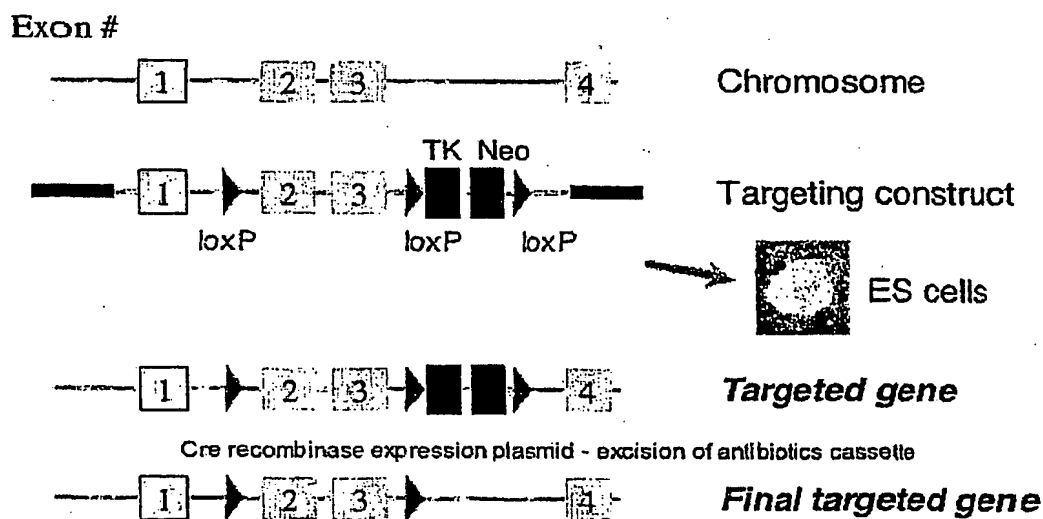
**LoxP site 3: Intron 3 5' of genomic XcmI site**  
underlined sequence = loxP site, cloning sites and partial Neo gene

35 gttttcat\*accaccgcgggtcccgggc\*gatat\*ttcaccttgtc\*ag\*cggtgttgtgtggtgtaaatg  
ttcgcgattgtttcgaaagcccc\*agcaccgcagtaagtcacggctcgggtacgtagacgatatcgctc  
gcgcgaacccagggccaccagcaagttgcgtggtggtggttttcccatcc\*gtggggac\*gtctatataa  
40 acc\*gcagtagcgtgggcattttctgctccgggcggaacttccgtggcttcttgcctgcccgcgagggcgcaa  
cgccgtacgtcgggttgcctatggccgcgagaacgcgcagcctgggtcgaacgcagacgcgtgtgtagggcgg  
ggtacgaagccatacgcgcttctacaaggcgctggccgaagaggtgcgggagtttcacgccaccaagatct  
gcggcacgctgttgacgctgttaagcgggtcgctgcagggtcgctcggtgttcgaggccacacgcgtcacc  
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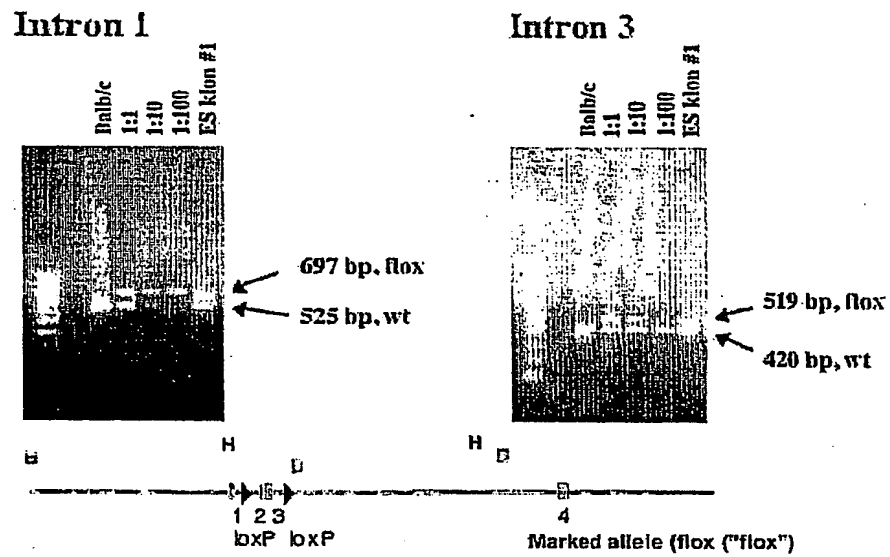
Fig. 4A Schematic representation of genetic manipulation.

### Serca2 (*atp2a2*) gene modification



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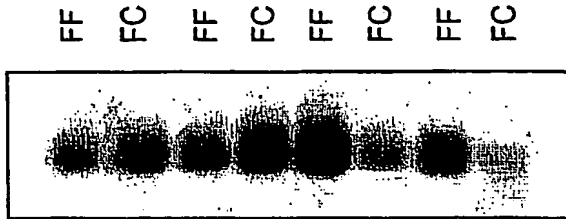
**Fig 4 B: Verification of Serca locus targeting events offspring from chimeric mice.**



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**Fig. 6 Cardiac ANP mRNA expression.**

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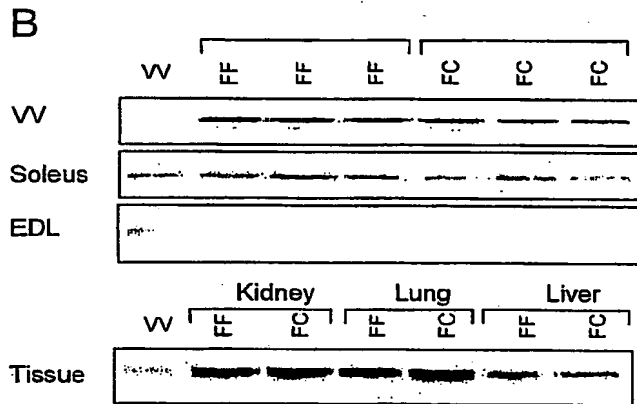
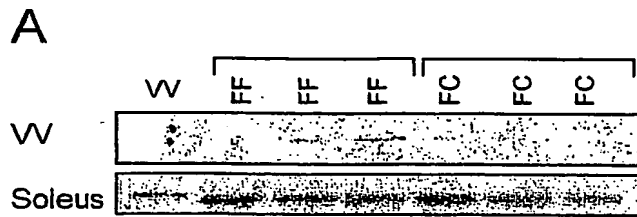


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Fig. 7 Serca2 protein expression.

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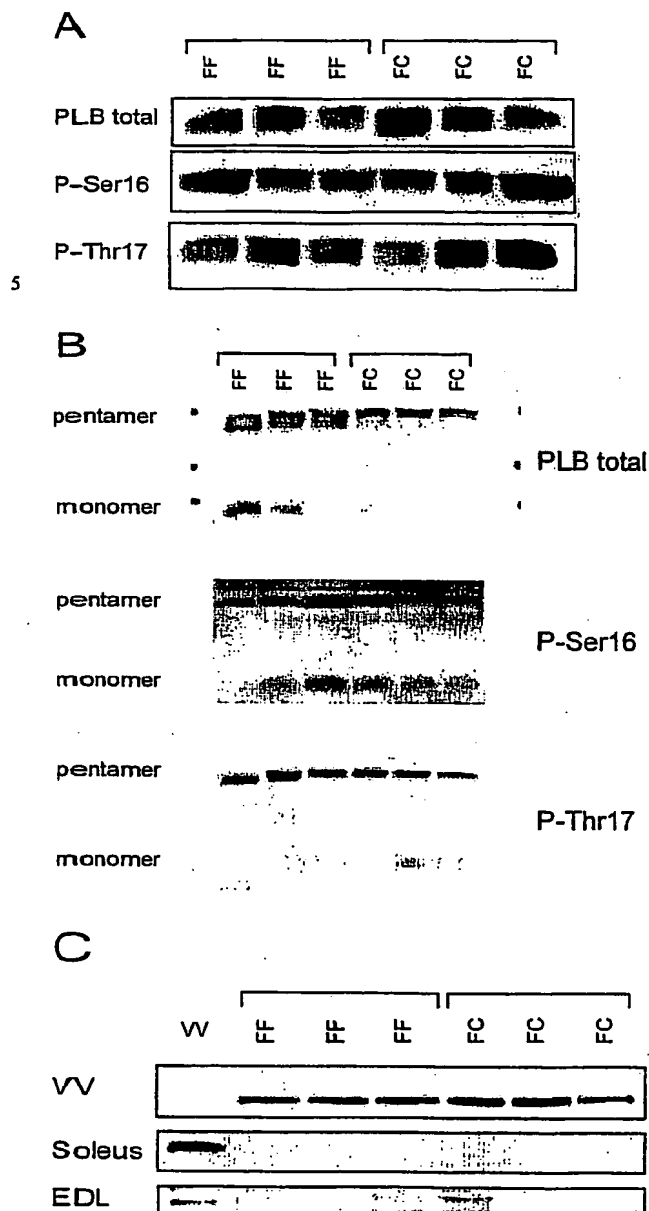


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**Fig. 8 Compensatory mechanisms in *Serca<sup>flox</sup>* MLC-2 $\nu$ -Cre mice.**



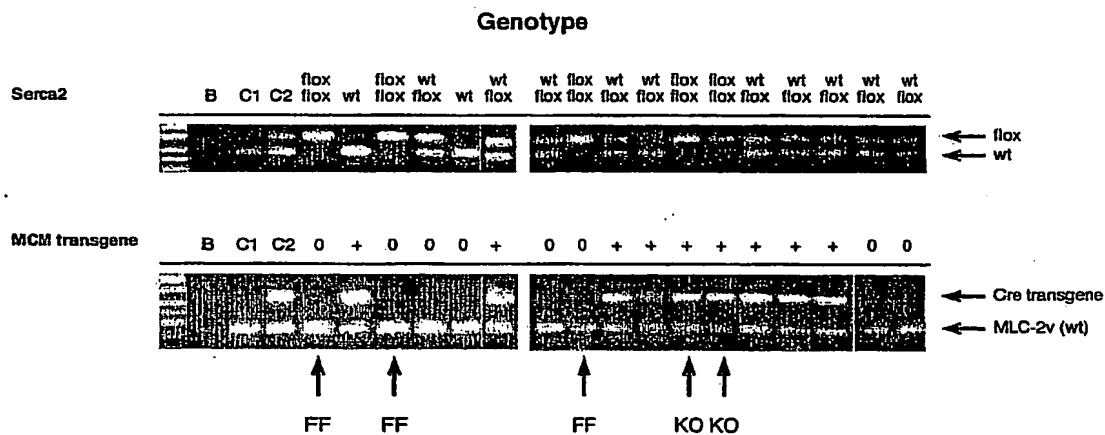
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DECT A1111 ADIE CDDV

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**Figure 9 Genotypes PCR**

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10 Generation of animals with Serca2<sup>flox</sup> and MCM transgene alleles.

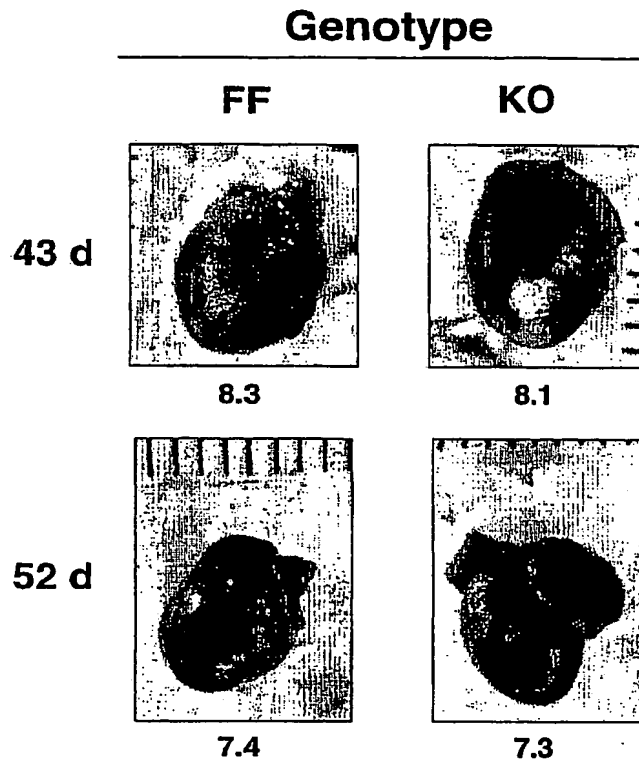
Genotypes FF. Serca2<sup>flox/flox</sup>; KO, Serca2<sup>flox/flox</sup> MCM

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Figure 10

Heart morphology

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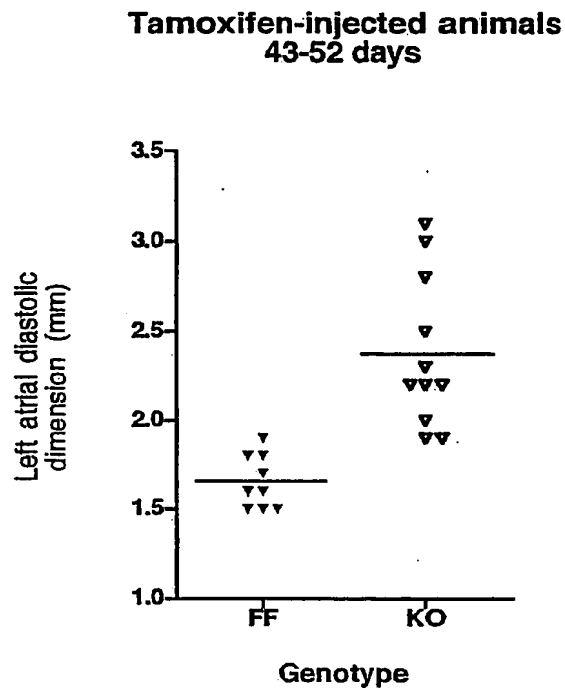
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**Figure 11** Pilot series left atrial diastolic diameter.

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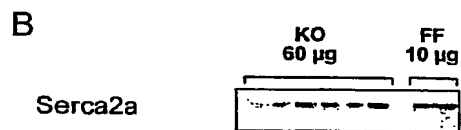
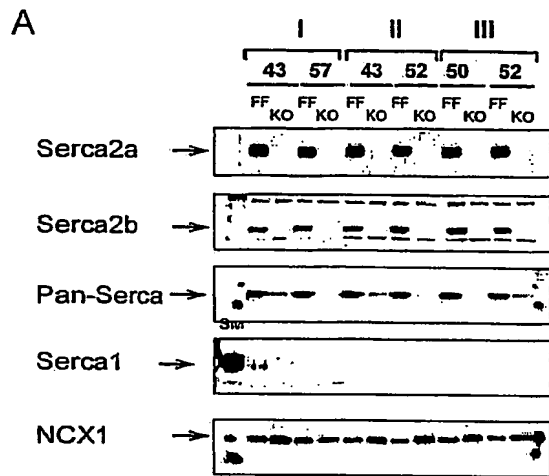
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**Figure 12 Serca protein content in tamoxifen-induced FF and KO mice**

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